SEQUENCE LISTING

<110> Sheppard, Paul O.
Baindur, Nand
Deisher, Theresa A.
Bishop, Paul D.

<120> DISINTEGRIN HOMOLOG

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cag caa aaa cat aat aag gct gtc cat ctg gcc cag gca agc ttc cag        95
Gln Gln Lys His Asn Lys Ala Val His Leu Ala Gln Ala Ser Phe Gln
                20                  25                  30

att gaa gcc ttc ggc tcc aaa ttc att ctt gac ctc ata ctg aac aat       143
Ile Glu Ala Phe Gly Ser Lys Phe Ile Leu Asp Leu Ile Leu Asn Asn
            35                  40                  45

ggt ttg ttg tct tct gat tat gtg gag att cac tac gaa aat ggg aaa       191
Gly Leu Leu Ser Ser Asp Tyr Val Glu Ile His Tyr Glu Asn Gly Lys
        50                  55                  60
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cca cag tac tct aag ggt gga gag cac tgt tac tac cat gga agc atc      239
Pro Gln Tyr Ser Lys Gly Gly Glu His Cys Tyr Tyr His Gly Ser Ile
     65                  70                  75

aga ggc gtc aaa gac tcc aag gtg gct ctg tca acc tgc aat gga ctt      287
Arg Gly Val Lys Asp Ser Lys Val Ala Leu Ser Thr Cys Asn Gly Leu
 80                  85                  90                  95

cat ggc atg ttt gaa gat gat acc ttc gtg tat atg ata gag cca cta      335
His Gly Met Phe Glu Asp Asp Thr Phe Val Tyr Met Ile Glu Pro Leu
                100                 105                 110

gag ctg gtt cat gat gag aaa agc aca ggt cga cca cat ata atc cag      383
Glu Leu Val His Asp Glu Lys Ser Thr Gly Arg Pro His Ile Ile Gln
            115                 120                 125

aaa acc ttg gca gga cag tat tct aag caa atg aag aat ctc act atg      431
Lys Thr Leu Ala Gly Gln Tyr Ser Lys Gln Met Lys Asn Leu Thr Met
        130                 135                 140

gaa aga ggt gac cag tgg ccc ttt ctc tct gaa tta cag tgg ttg aaa      479
Glu Arg Gly Asp Gln Trp Pro Phe Leu Ser Glu Leu Gln Trp Leu Lys
    145                 150                 155

aga agg aag aga gca gtg aat cca tca cgt ggt ata ttt gaa gaa atg      527
Arg Arg Lys Arg Ala Val Asn Pro Ser Arg Gly Ile Phe Glu Glu Met
160                 165                 170                 175

aaa tat ttg gaa ctt atg att ggt aat gat cac aaa acg tat aag aag      575
Lys Tyr Leu Glu Leu Met Ile Gly Asn Asp His Lys Thr Tyr Lys Lys
                180                 185                 190

cat cgc tct tct cat gca cat acc aac aac ttt gca aag tcc gtg gtc      623
His Arg Ser Ser His Ala His Thr Asn Asn Phe Ala Lys Ser Val Val
            195                 200                 205

aac ctt gtg gat tct att tac aag gag cag ctc aac acc agg gtt gtc      671
Asn Leu Val Asp Ser Ile Tyr Lys Glu Gln Leu Asn Thr Arg Val Val
        210                 215                 220

ctg gtg gct gta gag acc tgg act gag aag gat cag att gac atc acc      719
Leu Val Ala Val Glu Thr Trp Thr Glu Lys Asp Gln Ile Asp Ile Thr
    225                 230                 235
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acc aac cct gtg cag atg ctc cat gag ttc tca aaa tac cgg cag cgc      767
Thr Asn Pro Val Gln Met Leu His Glu Phe Ser Lys Tyr Arg Gln Arg
240                 245                 250                 255

att aag cag cat gct gat gct gtg cac ctc atc tcg cgg gtg aca ttt      815
Ile Lys Gln His Ala Asp Ala Val His Leu Ile Ser Arg Val Thr Phe
                260                 265                 270

cac tat aag aga agc agt ctg agt tac ttt gaa ggt gtc tgt tct cgc      863
His Tyr Lys Arg Ser Ser Leu Ser Tyr Phe Glu Gly Val Cys Ser Arg
            275                 280                 285

aca aga gga gtt ggt gtg aat gag tat ggt ctt cca atg gca gtg gca      911
Thr Arg Gly Val Gly Val Asn Glu Tyr Gly Leu Pro Met Ala Val Ala
        290                 295                 300

caa gta tta tcg cag agc ctg gct caa aac ctt gga atc caa tgg gaa      959
Gln Val Leu Ser Gln Ser Leu Ala Gln Asn Leu Gly Ile Gln Trp Glu
    305                 310                 315

cct tct agc aga aag cca aaa tgt gac tgc aca gaa tcc tgg ggt ggc     1007
Pro Ser Ser Arg Lys Pro Lys Cys Asp Cys Thr Glu Ser Trp Gly Gly
320                 325                 330                 335

tgc atc atg gag gaa aca ggg gtg tcc cat tct cga aaa ttt tca aag     1055
Cys Ile Met Glu Glu Thr Gly Val Ser His Ser Arg Lys Phe Ser Lys
                340                 345                 350

tgc agc att ttg gag tat aga gac ttt tta cag aga gga ggt gga gcc     1103
Cys Ser Ile Leu Glu Tyr Arg Asp Phe Leu Gln Arg Gly Gly Gly Ala
            355                 360                 365

tgc ctt ttc aac agg cca aca aag cta ttt gag ccc acg gaa tgt gga     1151
Cys Leu Phe Asn Arg Pro Thr Lys Leu Phe Glu Pro Thr Glu Cys Gly
        370                 375                 380

aat gga tac gtg gaa gct ggg gag gag tgt gat tgt ggt ttt cat gtg     1199
Asn Gly Tyr Val Glu Ala Gly Glu Glu Cys Asp Cys Gly Phe His Val
    385                 390                 395

gaa tgc tat gga tta tgc tgt aag aaa tgt tcc ctc tcc aac ggg gct     1247
Glu Cys Tyr Gly Leu Cys Cys Lys Lys Cys Ser Leu Ser Asn Gly Ala
400                 405                 410                 415
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cac tgc agc gac ggg ccc tgc tgt aac aat acc tca tgt ctt ttt cag      1295
His Cys Ser Asp Gly Pro Cys Cys Asn Asn Thr Ser Cys Leu Phe Gln
                420                 425                 430

cca cga ggg tat gaa tgc cgg gat gct gtg aac gag tgt gat att act      1343
Pro Arg Gly Tyr Glu Cys Arg Asp Ala Val Asn Glu Cys Asp Ile Thr
            435                 440                 445

gaa tat tgt act gga gac tct ggt cag tgc cca cca aat ctt cat aag      1391
Glu Tyr Cys Thr Gly Asp Ser Gly Gln Cys Pro Pro Asn Leu His Lys
        450                 455                 460

caa gac gga tat gca tgc aat caa aat cag ggc cgc tgc tac aat ggc      1439
Gln Asp Gly Tyr Ala Cys Asn Gln Asn Gln Gly Arg Cys Tyr Asn Gly
    465                 470                 475

gag tgc aag acc aga gac aac cag tgt cag tac atc tgg gga aca aag      1487
Glu Cys Lys Thr Arg Asp Asn Gln Cys Gln Tyr Ile Trp Gly Thr Lys
480                 485                 490                 495

gct gca ggg tct gac aag ttc tgc tat gaa aag ctg aat aca gaa ggc      1535
Ala Ala Gly Ser Asp Lys Phe Cys Tyr Glu Lys Leu Asn Thr Glu Gly
                500                 505                 510

act gag aag gga aac tgc ggg aag gat gga gac cgg tgg att cag tgc      1583
Thr Glu Lys Gly Asn Cys Gly Lys Asp Gly Asp Arg Trp Ile Gln Cys
            515                 520                 525

agc aaa cat gat gtg ttc tgt gga ttc tta ctc tgt acc aat ctt act      1631
Ser Lys His Asp Val Phe Cys Gly Phe Leu Leu Cys Thr Asn Leu Thr
        530                 535                 540

cga gct cca cgt att ggt caa ctt cag ggt gag atc att cca act tcc      1679
Arg Ala Pro Arg Ile Gly Gln Leu Gln Gly Glu Ile Ile Pro Thr Ser
    545                 550                 555

ttc tac cat caa ggc cgg gtg att gac tgc agt ggt gcc cat gta gtt      1727
Phe Tyr His Gln Gly Arg Val Ile Asp Cys Ser Gly Ala His Val Val
560                 565                 570                 575

tta gat gat gat acg gat gtg ggc tat gta gaa gat gga acg cca tgt      1775
Leu Asp Asp Asp Thr Asp Val Gly Tyr Val Glu Asp Gly Thr Pro Cys
                580                 585                 590
```

```
ggc ccg tct atg atg tgt tta gat cgg aag tgc cta caa att caa gcc     1823
Gly Pro Ser Met Met Cys Leu Asp Arg Lys Cys Leu Gln Ile Gln Ala
            595                 600                 605

cta aat atg agc agc tgt cca ctc gat tcc aag ggt aaa gtc tgt tcg     1871
Leu Asn Met Ser Ser Cys Pro Leu Asp Ser Lys Gly Lys Val Cys Ser
        610                 615                 620

ggc cat ggg gtg tgt agt aat gaa gcc acc tgc att tgt gat ttc acc     1919
Gly His Gly Val Cys Ser Asn Glu Ala Thr Cys Ile Cys Asp Phe Thr
    625                 630                 635

tgg gca ggg aca gat tgc agt atc cgg gat cca gtt agg aac ctt cac     1967
Trp Ala Gly Thr Asp Cys Ser Ile Arg Asp Pro Val Arg Asn Leu His
640                 645                 650                 655

ccc ccc aag gat gaa gga ccc aag ggt ttg tgt gat ttt ggt ttc aat     2015
Pro Pro Lys Asp Glu Gly Pro Lys Gly Leu Cys Asp Phe Gly Phe Asn
                660                 665                 670

tca tgg aat act gaa ttc gtt gac act gtt cca atg cac cag tat aac     2063
Ser Trp Asn Thr Glu Phe Val Asp Thr Val Pro Met His Gln Tyr Asn
            675                 680                 685

att cta att gac tta aga gga gac aca taagaatatc ngtttttgcc          2110
Ile Leu Ile Asp Leu Arg Gly Asp Thr
        690                 695

tttaaagtat ataatttatg ttactgccaa attaaggatt ctgatatatc atatttttaa   2170
aatgtgtttg aattacttct tagtctagaa ctgagattgg gaagaagtaa atatacacat   2230
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            20                  25                  30
Glu Ala Phe Gly Ser Lys Phe Ile Leu Asp Leu Ile Leu Asn Asn Gly
        35                  40                  45
```

```
Leu Leu Ser Ser Asp Tyr Val Glu Ile His Tyr Glu Asn Gly Lys Pro
    50                  55                  60
Gln Tyr Ser Lys Gly Gly Glu His Cys Tyr Tyr His Gly Ser Ile Arg
65                  70                  75                  80
Gly Val Lys Asp Ser Lys Val Ala Leu Ser Thr Cys Asn Gly Leu His
                85                  90                  95
Gly Met Phe Glu Asp Asp Thr Phe Val Tyr Met Ile Glu Pro Leu Glu
            100                 105                 110
Leu Val His Asp Glu Lys Ser Thr Gly Arg Pro His Ile Ile Gln Lys
        115                 120                 125
Thr Leu Ala Gly Gln Tyr Ser Lys Gln Met Lys Asn Leu Thr Met Glu
    130                 135                 140
Arg Gly Asp Gln Trp Pro Phe Leu Ser Glu Leu Gln Trp Leu Lys Arg
145                 150                 155                 160
Arg Lys Arg Ala Val Asn Pro Ser Arg Gly Ile Phe Glu Glu Met Lys
                165                 170                 175
Tyr Leu Glu Leu Met Ile Gly Asn Asp His Lys Thr Tyr Lys Lys His
            180                 185                 190
Arg Ser Ser His Ala His Thr Asn Asn Phe Ala Lys Ser Val Val Asn
        195                 200                 205
Leu Val Asp Ser Ile Tyr Lys Glu Gln Leu Asn Thr Arg Val Val Leu
    210                 215                 220
Val Ala Val Glu Thr Trp Thr Glu Lys Asp Gln Ile Asp Ile Thr Thr
225                 230                 235                 240
Asn Pro Val Gln Met Leu His Glu Phe Ser Lys Tyr Arg Gln Arg Ile
                245                 250                 255
Lys Gln His Ala Asp Ala Val His Leu Ile Ser Arg Val Thr Phe His
            260                 265                 270
Tyr Lys Arg Ser Ser Leu Ser Tyr Phe Glu Gly Val Cys Ser Arg Thr
        275                 280                 285
Arg Gly Val Gly Val Asn Glu Tyr Gly Leu Pro Met Ala Val Ala Gln
    290                 295                 300
Val Leu Ser Gln Ser Leu Ala Gln Asn Leu Gly Ile Gln Trp Glu Pro
305                 310                 315                 320
Ser Ser Arg Lys Pro Lys Cys Asp Cys Thr Glu Ser Trp Gly Gly Cys
                325                 330                 335
Ile Met Glu Glu Thr Gly Val Ser His Ser Arg Lys Phe Ser Lys Cys
            340                 345                 350
Ser Ile Leu Glu Tyr Arg Asp Phe Leu Gln Arg Gly Gly Gly Ala Cys
        355                 360                 365
Leu Phe Asn Arg Pro Thr Lys Leu Phe Glu Pro Thr Glu Cys Gly Asn
    370                 375                 380
Gly Tyr Val Glu Ala Gly Glu Glu Cys Asp Cys Gly Phe His Val Glu
385                 390                 395                 400
```

```
Cys Tyr Gly Leu Cys Cys Lys Lys Cys Ser Leu Ser Asn Gly Ala His
                405                 410                 415
Cys Ser Asp Gly Pro Cys Cys Asn Asn Thr Ser Cys Leu Phe Gln Pro
            420                 425                 430
Arg Gly Tyr Glu Cys Arg Asp Ala Val Asn Glu Cys Asp Ile Thr Glu
        435                 440                 445
Tyr Cys Thr Gly Asp Ser Gly Gln Cys Pro Pro Asn Leu His Lys Gln
    450                 455                 460
Asp Gly Tyr Ala Cys Asn Gln Asn Gln Gly Arg Cys Tyr Asn Gly Glu
465                 470                 475                 480
Cys Lys Thr Arg Asp Asn Gln Cys Gln Tyr Ile Trp Gly Thr Lys Ala
                485                 490                 495
Ala Gly Ser Asp Lys Phe Cys Tyr Glu Lys Leu Asn Thr Glu Gly Thr
            500                 505                 510
Glu Lys Gly Asn Cys Gly Lys Asp Gly Asp Arg Trp Ile Gln Cys Ser
        515                 520                 525
Lys His Asp Val Phe Cys Gly Phe Leu Leu Cys Thr Asn Leu Thr Arg
    530                 535                 540
Ala Pro Arg Ile Gly Gln Leu Gln Gly Glu Ile Ile Pro Thr Ser Phe
545                 550                 555                 560
Tyr His Gln Gly Arg Val Ile Asp Cys Ser Gly Ala His Val Val Leu
                565                 570                 575
Asp Asp Asp Thr Asp Val Gly Tyr Val Glu Asp Gly Thr Pro Cys Gly
            580                 585                 590
Pro Ser Met Met Cys Leu Asp Arg Lys Cys Leu Gln Ile Gln Ala Leu
        595                 600                 605
Asn Met Ser Ser Cys Pro Leu Asp Ser Lys Gly Lys Val Cys Ser Gly
    610                 615                 620
His Gly Val Cys Ser Asn Glu Ala Thr Cys Ile Cys Asp Phe Thr Trp
625                 630                 635                 640
Ala Gly Thr Asp Cys Ser Ile Arg Asp Pro Val Arg Asn Leu His Pro
                645                 650                 655
Pro Lys Asp Glu Gly Pro Lys Gly Leu Cys Asp Phe Gly Phe Asn Ser
            660                 665                 670
Trp Asn Thr Glu Phe Val Asp Thr Val Pro Met His Gln Tyr Asn Ile
        675                 680                 685
Leu Ile Asp Leu Arg Gly Asp Thr
    690                 695
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aartgyytnc arathcargc nytnaayatg wsnwsntgyc cnytngayws naarggnaar 1860
gtntgywsng gncayggngt ntgywsnaay gargcnacnt gyathtgyga yttyacntgg 1920
gcnggnacng aytgywsnat hmgngayccn gtnmgnaayy tncayccncc naargaygar 1980
ggnccnaarg gnytntgyga yttyggntty aaywsntgga ayacngartt ygtngayacn 2040

gtnccnatgc aycartayaa yathytnath gayytnmgng gngayacn 2088

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